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99; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    29 AVTONIME - LVECAS AISSAUPPSSSONSFVP----- EQRECEGYLRNPSL 74
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Best Local Similarity 13.3%, Fred No. 0.0025,
Matches 27, Conservative 8; Mismatches 28, Indels
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/632,511A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 2, Application US/08632511A
Patent No. 621850A
GENERAL INFORMATION:
APPLICANT: Kraft Anti-microbial proteins
NUMBER OF ENDERNICH, Anti-microbial proteins
NUMBER OF SEQUENCES: 9
CORPESSONDENCE APPROX
ADDRESSEE: SANDOZ ATPO INC
STREET: Palo Alto
STREET: California Avenue
CITY: Palo Alto
STATE: California
COMPUTY: USA
ZIP: CA94304-1104
COMPUTER: FLODEY disk
COMPUTER: FLODEY COMPATIBLE
COMPUTER: FLODEY DISC
COMPUTER: FLODEY COMPATIBLE
COMPUTER: FLODEY COMPATIBLE
COMPUTER: FLODEY COMPATIBLE
COMPUTER: FLODEY COMPATIBLE
COMPATING SYSTEM: PCLODE/MS-DOS
COMPUTER: FLODEY COMPATIBLE
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ATTORNEY, AGENT INFORMATION.
NAME: MATCHAL WYDEN, LDYNN
REGISTRATION NUMBER: 34,869
FEFFENCE/DOCKET NUMBER: 137-
TELECHMULICATION INFORMATION
TELEPHONE: 650,354,358
INFORMATION FOR SEQ. ID NO. 2-
SEQUENCE CHARACTERISTICS:
PEGISTPATION NUMBER: 33,367
REPEBENCE, POOPET UNMBER:
TELECOMMUNICATION
TELEPHONE: 202-371-8850
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         34,869
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    57 AGAYNAGPAAGIPSPÖÖVSVP 77
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                                                                                                                                                                                                                                                                                                            90 amino acids
                                                                                                                                                                                                                                INFORMATION FOR SEG ID NO. SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 91 amino ac.ds
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MOLEGULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                single
                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear
MOLECULE TYPE: peptide
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                                                                                                                                                                                                                                                                                                                                                        amino acid
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HYPOTHETICAL: N
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US-09-632-511A-2
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                                                                                                                                                                                                                                                                                                                LENGIH:
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12; Gaps
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                                                                                                                        Query Match
17.0%; Score 87; DB 4; Length 91;
Best Local Similarity 24.4%; Pred R1.....333;
Matches 21; Conservative 14; Mishatches 31; Indelic
Query Match
Best Local Similarity 26.9%, Fred. No. 0.0033,
Matches 21, Conservative 14, Mismatches 31, Indels
                                                                                                                                                                                                                                                                                                                                              Sequence 7, Application US/09C91590A
FRIEND 10. 6242574
GRBEAL INFORMATION. Klaus
APPLICANT: NetSen, Klaus
APPLICANT: Kroll Kristensen, Anne
APPLICANT: Kroll Kristensen, Anne
TITLE OF INVENTOR 10 Jane
TILE FEFFENTS: 3.13 1111/NA.A.(3532/FCT
CURRENT AFFLICATION NYMEEF 1999-65.0
FRIOR FILING DATE: 1996-12-20
PRIOR FILING DATE: 1996-12-20
PRIOR FILING DATE: 1996-12-21
PRIOR FILING DATE: 1996-12-22
PRIOR FILING DATE: 1996-12-22
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APPLICANT: Kroll Kristensen, Anne
APPLICANT: Kroll Kristensen, Anne
TITLE: VERONT: Brunstedt, Janne
TITLE: VERNETISM
ANTA: MINISTERN
CUPPENT AFFLICATI: NYMBER: US/03/091,590A
CUPPENT AFFLICATI: NYMBER: US/03/091,590A
CUPPENT AFFLICATI: NYMBER: US/03/091,590A
CUPPENT AFFLICATI: NYMBER: US/03/091,590A
PRIOR PRILING DATE: 1996-12-0
PRIOR FILING DATE: 1996-12-0
PRIOR FILING NAMER: 36 9526238.2
PRIOR FILING NAMER: 36 9526238.2
PRIOR FILING NAMER: 35 9526238.2
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                                                                                                                                                                                     78 VSPAAAQKIASQCGVSIP 95
                                                                                                                                                                                                                                  61 INYGKAASLPROCGUSIP 78
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             61 INYGKAASLPROCGVSIP 78
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT
ORGANISM: Beta vulgaris
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: PRT ORGANISM: Beta vulgaris
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       , Patent No. 6242574
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                       RESULT 10.
US-09-091-590A-7
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